Ensemble Learning

Nipun Batra and teaching staff

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IIT Gandhinagar

Use multiple models for prediction.

Most winning entries of Kaggle competition using ensemble learning.

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Example:

Classifier 1 - Good

Classifier 2 - Good

Classifier 3 - Bad

Using Majority Voting, we predict Good.

Use multiple models for prediction.

Most winning entries of Kaggle competition using ensemble learning.

Example:

Regressor 1 - 20

Regressor 2 - 30

Regressor 3 - 30

Using Average, we predict $\frac{80}{3}$

Based on Ensemble methods in ML by Dietterich

Three reasons why ensembles make sense:

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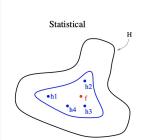
1) Statistical: Sometimes if data is less, many competing hypothesis can be learnt all giving same accuracy on training data.

Based on Ensemble methods in ML by Dietterich

Three reasons why ensembles make sense:

1) Statistical: Sometimes if data is less, many competing hypothesis can be learnt all giving same accuracy on training data.

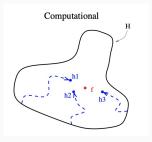
Eg. We can learn many decision trees for the same data giving same accuracy.



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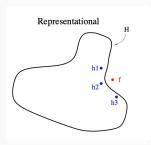
Eg. Decision Trees employ greedy critera



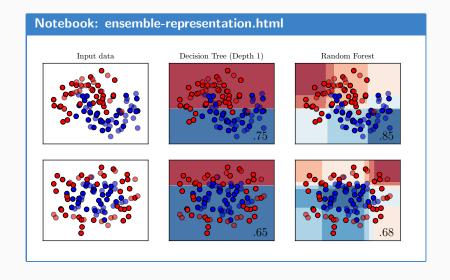
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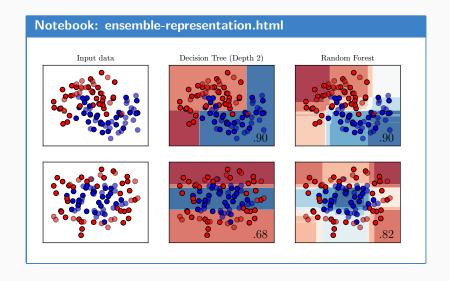
Eg. Decision Trees can only learn axis-parallel splits.



Representation of Limited Depth DTs vs RFs



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- 3) Two classifiers are diverse: if they make different errors on new data points

Imagine that we have an ensemble of three classifiers (h_1, h_2, h_3) and consider a new case x.

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If the three classifiers are identical, i.e. not diverse, then when $h_1(x)$ is wrong $h_2(x)$ and $h_3(x)$ will also be wrong.

However, if the errors made by the classifiers are uncorrelated, then when $h_1(x)$ is wrong, $h_2(x)$ and $h_3(x)$ may be correct, so that a majority vote will correctly class.

Intuition for Ensemble Methods from Quantitative Perspective

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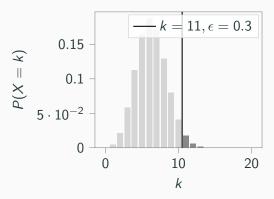
Error Probability of each model $= \varepsilon = 0.3$

Pr(ensemble being wrong) =
$${}^3C_2(\varepsilon^2)(1-\varepsilon)^{3-2}+{}^3C_3(\varepsilon^3)(1-\varepsilon)^{3-3}$$

= $0.19 \le 0.3$

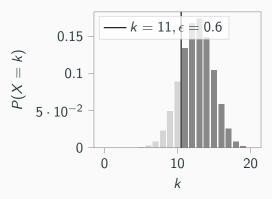
Some calculations

Probability that majority vote (11 out of 21) is wrong = 0.026



Some calculations

Probability that majority vote (11 out of 21) is wrong = 0.826



Where does ensemble learning not work well?

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- The base model is bad.
- All models give similar prediction or the models are highly correlated.

Also known as Bootstrap Aggregation.

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Key idea: Reduce Variance

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How to learn different classifiers while feeding in the same data?

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Think about cross-validation!

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Key idea: Reduce Variance

How to learn different classifiers while feeding in the same data?

Think about cross-validation!

We will create multiple datasets from our single dataset using "sampling with replacement".

Consider our dataset has n samples, $D_1, D_2, D_3, \ldots, D_n$. For each model in the ensemble, we create a new dataset of size n by sampling uniformly with replacement.

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Round 1 : $D_1, D_3, D_6, D_1, \dots, D_n$ Round 2 : $D_2, D_4, D_1, D_{80}, \dots, D_3$:

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$$D_1, D_3, D_6, D_1, \dots, D_n$$

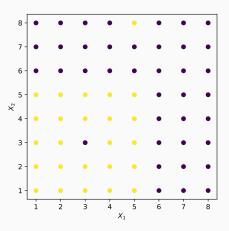
Round 2:
$$D_2, D_4, D_1, D_{80}, \dots, D_3$$

:

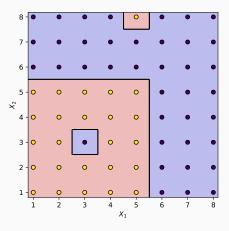
Repetition of samples is possible.

We can train the same classifier/models on each of these different "Bagging Rounds".

Consider the dataset below. Points (3,3) and (5,8) are anomalies.

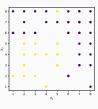


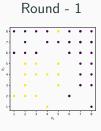
Decision Boundary for decision tree with depth 6.

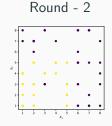


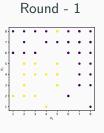
Lets use bagging with ensemble of 5 trees.

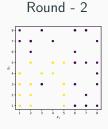
Round - 1

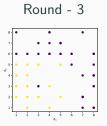


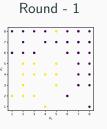




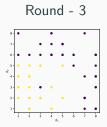


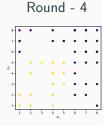


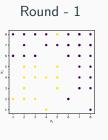


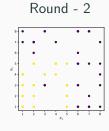


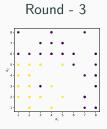


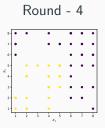


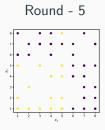




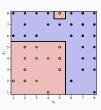






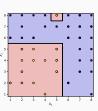


Round - 1



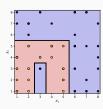
 $\mathsf{Tree}\ \mathsf{Depth} = \mathsf{4}$



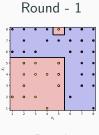


Tree Depth = 4

Round - 2

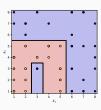


 $\mathsf{Tree}\ \mathsf{Depth} = \mathsf{5}$



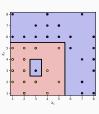
Tree Depth = 4



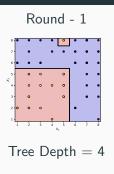


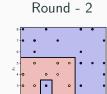
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Round - 3

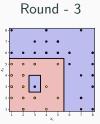


Tree Depth = 5



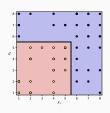


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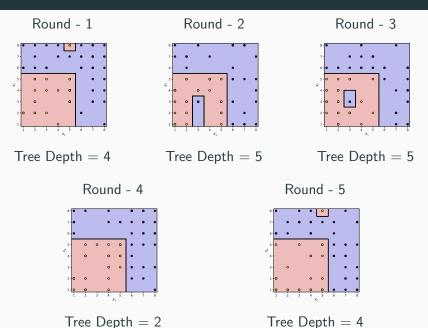


Tree Depth = 5

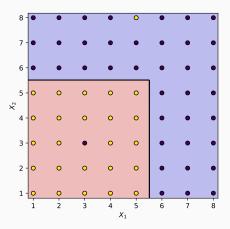




Tree Depth = 2



Using majority voting to combine all predictions, we get the decision boundary below.



Summary

- We take "strong" learners and combine them to reduce variance.
- All learners are independent of each other.

Boosting

• We take "weak" learners and combine them to reduce bias.

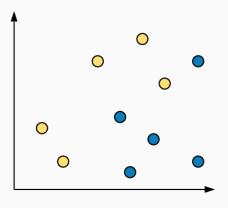
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- Incremental building: Incrementally try to classify "harder" samples correctly.

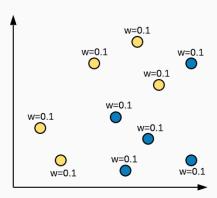
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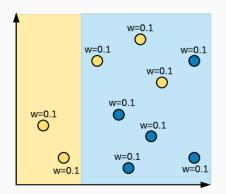
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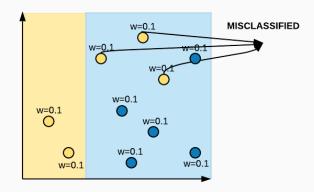
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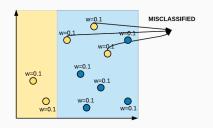
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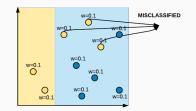
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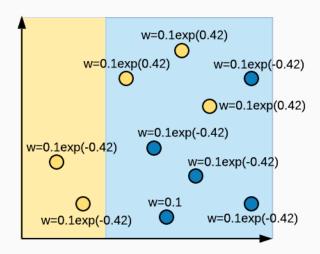
2.3 Compute
$$\alpha_m = \frac{1}{2} log_e \left(\frac{1 - err_m}{err_m} \right)$$

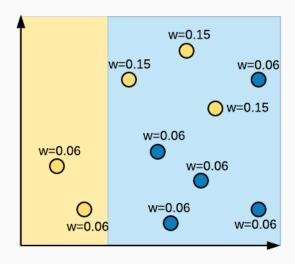


$$err_1 = \frac{0.3}{1}$$
 $\alpha_1 = \frac{1}{2}log\left(\frac{1-0.3}{0.3}\right) = 0.42$

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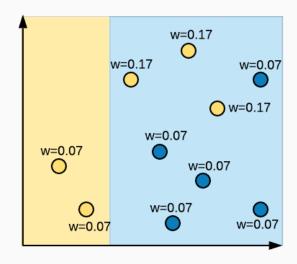


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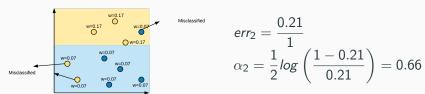


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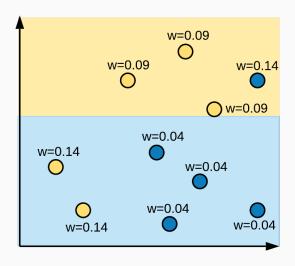
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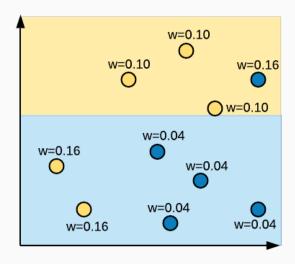
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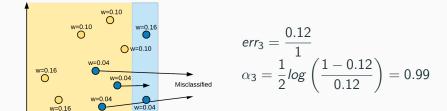
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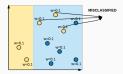
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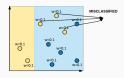
Intuitively, after each iteration, importance of wrongly classified samples is increased by increasing their weights and importance of correctly classified samples is decreased by decreasing their weights.

Testing

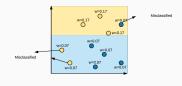
- For each sample x, compute the prediction of each classifier h_m(x).
- Final prediction is the sign of the sum of weighted predictions, given as:
- SIGN $(\alpha_1 h_1(x) + \alpha_2 h_2(x) + ... + \alpha_M h_M(x))$



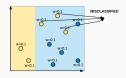
$$\alpha_1 = 0.42$$



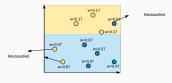
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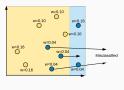
$$\alpha_2 = 0.66$$



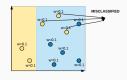
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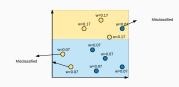
$$\alpha_2 = 0.66$$



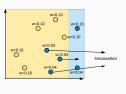
$$\alpha_3 = 0.99$$



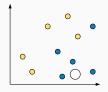
$$\alpha_1 = 0.42$$



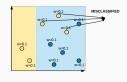
$$\alpha_2 = 0.66$$



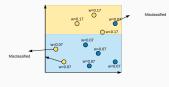
$$\alpha_3 = 0.99$$



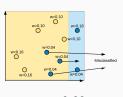
Example



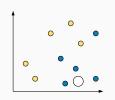
$$\alpha_1 = 0.42$$



$$\alpha_2 = 0.66$$

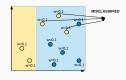


 $\alpha_3 = 0.99$

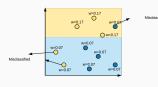


Let us say, yellow class is +1 and blue class is -1

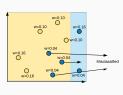
Example



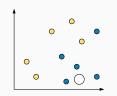
$$\alpha_1 = 0.42$$



$$\alpha_2 = 0.66$$



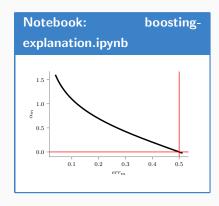
$$\alpha_3 = 0.99$$



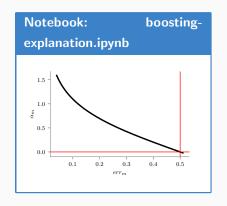
Let us say, yellow class is +1 and blue class is -1Prediction = SIGN(0.42*-1 + 0.66*-1 + 0.99*+1) = Negative = blue

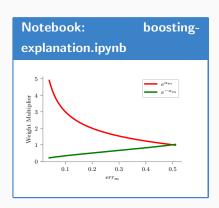
Intuition behind weight update formula

Intuition behind weight update formula



Intuition behind weight update formula





ADABoost for regresion

From Paper: Improving Regressors using Boosting Techniques

Our problem will be that the modeling error is also nonzero because we have to determine the model in the presence of noise. Since we don't know the probability distributions, we approximate the expectation of the ME and PE using the sample ME (if the truth is known) and sample PE and then average over multiple experiments.

In the following discussion, we detail both bagging and boosting. We then discuss how to build trees which are the basic building blocks of our regression machines and use these ensembles on some standard test functions.

2. BAGGING

The following is a paraphrase of Breiman (1996b) with some difference in notation. Suppose we pick with replacement N_1 examples from the training set of size N_{\perp} and call the k'th set of observations O_{\perp} . Based on these observations, we form a predictor $v^{(p)}(\mathbf{x}, O_k)$. Because we are sampling with replacement, we may have multiple observations or no observations of a particular training example. Sampling with replacement is sometimes termed bootstrap sampling [Efron and Tibshirani (1993)] and therefore this method is called bootstrap aggregating or bagging for short. The ensemble predictor is formed from the approximation to the expectation over all the observation sets, i.e. $E_O[y^{(p)}(\mathbf{x}, O)]$ by using the average of the outputs of all the predictors. Breiman discusses which algorithms are good candidates for predictors and concludes that the best predictors are unstable, i.e., a small change in the training set O causes a large change in the predictor $v^{(p)}(\mathbf{x}, O_k)$. Good candidates are regression trees and neural nets.

3. BOOSTING

In bagging, each training example is equally likely to be picked. In boosting, the probability of a particular example being in the training set of a particular machine depends on the performance of the prior machines on that example. The following is a modification of Adaboost.R [Freund and Schapire (1996a)].

Initially, to each training pattern we assign a weight $w_i=1$ $i=1,...,N_1$

Repeat the following while the average loss \bar{L} defined

set. Each machine makes a hypothesis: h.:x→v

 Pass every member of the training set through this machine to obtain a prediction y_i^(p)(x_i) i=1,...N₁.

4. Calculate a loss for each training sample $L_l = L \left[|y_l^{(p)}(\mathbf{x}_l) - y_l| \right]$. The loss L may be of any functional form as long as $L \in [0,1]$. If we let

$$D=sup \mid y_i^{(p)}(\mathbf{x}_i) - y_i \mid i=1,...,N_1$$

then we have three candidate loss functions:

$$L_i = \frac{|y_i^{(p)}(x_i) - y_i|}{D}$$
 (linear)

$$L_i = \frac{\mid y_i^{(p)}(\pmb{x}_i) - y_i \mid^2}{D^2} \qquad (square~law)$$

$$L_{i}{=}1-exp\left\{ \frac{-\mid y_{i}^{(p)}(\boldsymbol{x}_{i})-y_{i}\mid}{D}\right\} \qquad (exponential)$$

5. Calculate an average loss: $\overline{L} = \sum_{i=1}^{N_1} L_i p_i$

6. Form $\beta = \frac{\overline{L}}{1-\overline{L}}$, β is a measure of confidence in the predictor. Low β means high confidence in the prediction.

7. Update the weights: w₁→w₁β**![1-L₁], where ** indicates exponentiation. The smaller the loss, the more the weight is reduced making the probability smaller that this pattern will be picked as a member of the training set for the next machine in the ensemble.

 For a particular input x_i, each of the T machines makes a prediction h_i, t=1,...,T. Obtain the cumulative prediction h_i using the T predictors:

Random Forest

- Random Forest is an ensemble of decision trees.
- We have two types of bagging: bootstrap (on data) and random subspace (of features).
- As features are randomly selected, we learn decorrelated trees and helps in reducing variance.

Random Forest

There are 3 parameters while training a random forest number of trees, number of features (m), maximum depth.

Training Algorithm

- For i^{th} tree $(i \in \{1 \cdots N\})$, select n samples from total N samples with replacement.
- Learn Decision Tree on selected samples for i^{th} round.

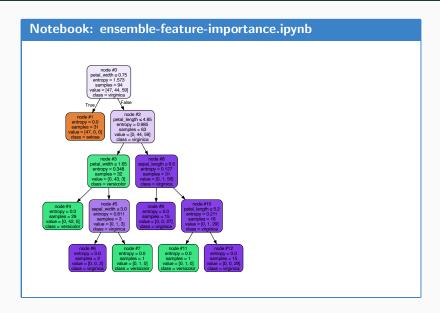
Learning Decision Tree (for RF)

 For each split, select m features from total available M features and train a decision tree on selected features

Dataset

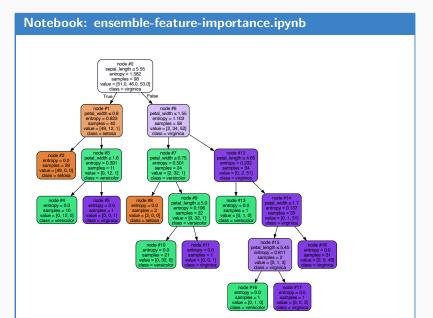
sepal_length	sepal_width	petal_length	petal_width	species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
6.7	3.0	5.2	2.3	virginica
6.3	2.5	5.0	1.9	virginica
6.5	3.0	5.2	2.0	virginica
6.2	3.4	5.4	2.3	virginica
5.9	3.0	5.1	1.8	virginica
	5.1 4.9 4.7 4.6 5.0 6.7 6.3 6.5 6.2	5.1 3.5 4.9 3.0 4.7 3.2 4.6 3.1 5.0 3.6 6.7 3.0 6.3 2.5 6.5 3.0 6.2 3.4	5.1 3.5 1.4 4.9 3.0 1.4 4.7 3.2 1.3 4.6 3.1 1.5 5.0 3.6 1.4 6.7 3.0 5.2 6.3 2.5 5.0 6.5 3.0 5.2 6.2 3.4 5.4	5.1 3.5 1.4 0.2 4.9 3.0 1.4 0.2 4.7 3.2 1.3 0.2 4.6 3.1 1.5 0.2 5.0 3.6 1.4 0.2 6.7 3.0 5.2 2.3 6.3 2.5 5.0 1.9 6.5 3.0 5.2 2.0 6.2 3.4 5.4 2.3

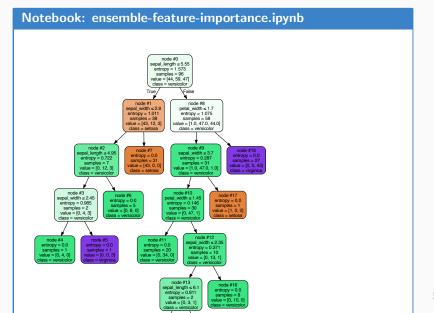
150 rows × 5 columns



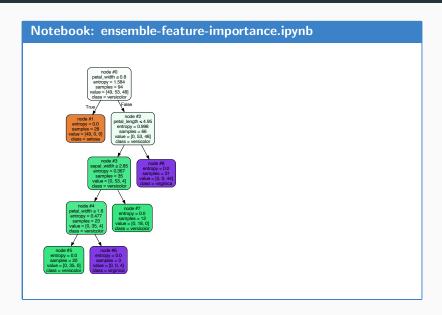
class = versicolor

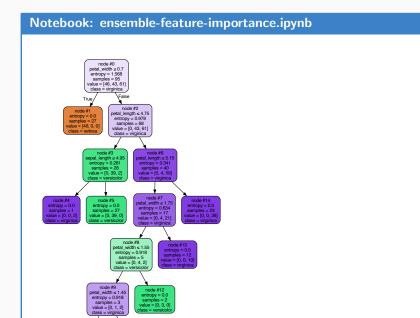
Notebook: ensemble-feature-importance.ipynb node #0 petal_width ≤ 0.8 entropy = 1.564samples = 100 value = [46.0, 62.0, 42.0] class = versicolor True node #2 node #1 petal width ≤ 1.75 entropy = 0.0entropy = 0.973 samples = 33 samples = 67 value = [46, 0, 0] value = [0, 62, 42] class = setosa class = versicolor node #3 petal_length ≤ 4.95 oetal_length ≤ 4.85 entropy = 0.387 samples = 39 samples = 28 value = [0, 61, 5] value = [0, 1, 37] class = versicolor class = virginica node #11 node #5 node #4 node #14 petal_length ≤ 5.45 sepal_width ≤ 3.1 entropy = 0.0entropy = 0.954 entropy = 0.918samples = 35 samples = 26 samples = 4 samples = 2 value = [0, 58, 0] value = [0, 0, 35] value = [0, 1, 2] value = [0, 3, 5] class = versicolo class = virginica class = virginica node #6 node #13 sepal width ≤ 2.45 entropy = 0.0entropy = 0.971 samples = 1 samples = 1 samples = 1 samples = 3 value = [0, 0, 3] value = [0, 0, 2]value = [0, 1, 0] value = [0, 3, 2]class = versicolo class = versicolor node #7 node #8 entropy = 0.0samples = 1 samples = 2 value = [0, 3, 0]





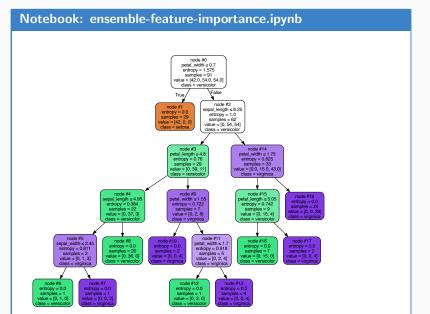
Notebook: ensemble-feature-importance.ipynb node #0 petal_width ≤ 0.7 entropy = 1.561 samples = 95 value = [50, 61, 39] class = versicolor node #2 node #1 petal width ≤ 1.75 entropy = 0.0entropy = 0.965 samples = 31 samples = 64 value = [50, 0, 0] value = [0, 61, 39] class = setosa class = versicolor node #3 node #10 petal_length ≤ 4.95 entropy = 0.435 samples = 41 value = [0, 61, 6] class = virginica class = versicolor node #5 node #4 petal_width ≤ 1.55 entropy = 0.0entropy = 0.994 samples = 35 samples = 6 value = [0, 56, 0] value = [0, 5, 6] lass = versicolor class = virginica node #7 petal length ≤ 5.45 entropy = 0.65 samples = 3 value = [0, 5, 1] class = versicolor node #8 entropy = 0.0samples = 2 samples = 1 value = [0, 5, 0] value = [0, 0, 1] dass = versicolo





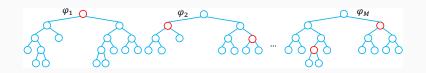
Notebook: ensemble-feature-importance.ipynb





Notebook: ensemble-feature-importance.ipynb node #0 petal_length ≤ 2.6 entropy = 1.57 samples = 96 value = [55.0, 40.0, 55.0] class = setosa True node #2 node #1 petal length ≤ 4.95 entropy = 0.0entropy = 0.982 samples = 36 samples = 60 value = [55, 0, 0] value = [0, 40, 55] class = setosa class = virginica node #3 sepal_length ≤ 5.95 entropy = 0.446 samples = 30 samples = 30 value = [0, 39, 4] value = [0, 1, 51] class = versicolor class = virginica node #5 node #4 node #9 petal_width ≤ 1.65 sepal_length ≤ 6.75 entropy = 0.0 entropy = 0.297 samples = 14 entropy = 0.722samples = 15 samples = 16 samples = 15 value = [0, 23, 0] value = [0, 16, 4] value = [0, 1, 18] lass = versicolor class = versicolor class = virginica node #11 node #6 petal width ≤ 2.0 entropy = 0.0entropy = 0.722 samples = 12 samples = 3 samples = 4 value = [0, 16, 0] value = [0, 1, 4] lass = versicolor class = virginica node #12 node #13 entropy = 0.0samples = 1 samples = 3 value = [0, 1, 0] value = [0, 0, 4] dass = versicolor class = virgini

Feature Importance¹



Importance of variable X_j for an ensemble of M trees φ_m is:

$$Imp(X_j) = \frac{1}{M} \sum_{m=1}^{M} \sum_{t \in \varphi_m} 1(j_t = j) \Big[p(t) \Delta i(t) \Big],$$

where j_t denotes the variable used at node t, $p(t) = N_t/N$ and $\Delta i(t)$ is the impurity reduction at node t:

$$\Delta i(t) = i(t) - \frac{N_{t_L}}{N_t}i(t_L) - \frac{N_{t_r}}{N_t}i(t_R)$$

¹Slide Courtesy Gilles Louppe

Computed Feature Importance

